

SEQUENCE LISTING

<110> Vogeli, Gabriel
Wood, Linda S.
Merchant, Kalpana

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 His Thr Arg Pro Trp Thr Ser Asn Thr Val Phe Leu Val Ser Leu Val
 85 90 95
 Ala Ala Asp Phe Leu Leu Ile Ser Asn Leu Pro Leu Arg Val Asp Tyr
 100 105 110
 Tyr Leu Leu His Glu Thr Trp Arg Phe Gly Ala Ala Ala Cys Lys Val
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 Asn Leu Phe Met Leu Ser Thr Asn Arg Thr Ala Ser Val Val Phe Leu
 130 135 140
 Thr Ala Ile Ala Leu Asn Arg Tyr Leu Lys Val Val Gln Pro His His
 145 150 155 160
 Val Leu Ser Arg Ala Ser Val Gly Ala Ala Ala Arg Val Ala Gly Gly
 165 170 175
 Leu Trp Val Gly Ile Leu Leu Leu Asn Gly His Leu Leu Leu Ser Thr
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 Phe Ser Gly Pro Ser Cys Leu Ser Tyr Arg Val Gly Thr Lys Pro Ser
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 Ala Ser Leu Arg Trp His Gln Ala Leu Tyr Leu Leu Glu Phe Phe Leu
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 Pro Leu Ala Leu Ile Leu Phe Ala Ile Val Ser Ile Gly Leu Thr Ile
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 Arg Asn Arg Gly Leu Gly Gly Gln Ala Gly Pro Gln Arg Ala Met Arg
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 Val Leu Ala Met Val Val Ala Val Tyr Thr Ile Cys Phe Leu Pro Ser
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Thr Tyr Leu Asn Ser Val Leu Asp Pro Val Leu Tyr Cys Phe Ser Ser
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Pro Asn Phe Leu His Gln Ser Arg Ala Leu Leu Gly Leu Thr Arg Gly
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Arg Gln Gly Pro Val Ser Asp Glu Ser Ser Tyr Gln Pro Ser Arg Gln
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Met Asn Thr Thr Val Met Gln Gly Phe
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Asn Arg Ser Glu Arg Cys Pro Arg Asp Thr Arg Ile Val Gln Leu Val
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Phe Pro Ala Leu Tyr Thr Val Val Phe Leu Thr Gly Ile Leu Leu Asn
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Thr Leu Ala Leu Trp Val Phe Val His Ile Pro Ser Ser Ser Thr Phe
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Ile Ile Tyr Leu Lys Asn Thr Leu Val Ala Asp Leu Ile Met Thr Leu
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Met Leu Pro Phe Lys Ile Leu Ser Asp Ser His Leu Ala Pro Trp Gln
75 80 85
ctc aga gct ttt gtg tgt cgt ttt tct tcg gtg ata ttt tat gag acc 460
Leu Arg Ala Phe Val Cys Arg Phe Ser Ser Val Ile Phe Tyr Glu Thr
90 95 100 105
atg tat gtg ggc atc gtg ctg tta ggg ctc ata gcc ttt gac aga ttc 508
Met Tyr Val Gly Ile Val Leu Leu Gly Leu Ile Ala Phe Asp Arg Phe
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Val His Ile Pro Ser Ser Ser Thr Phe Ile Ile Tyr Leu Lys Asn Thr
50 55 60
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Ser Asp Ser His Leu Ala Pro Trp Gln Leu Arg Ala Phe Val Cys Arg
85 90 95
Phe Ser Ser Val Ile Phe Tyr Glu Thr Met Tyr Val Gly Ile Val Leu
100 105 110
Leu Gly Leu Ile Ala Phe Asp Arg Phe Leu Lys Ile Ile Arg Pro Leu
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Arg Asn Ile Phe Leu Lys Lys Pro Val Phe Ala Lys Thr Val Ser Ile
130 135 140
Phe Ile Trp Val Phe Leu Val Phe Ile Ser Leu Pro Asn Met Ile Leu
145 150 155 160
Ser Asn Lys Glu Ala Thr Pro Ser Ser Val Lys Lys Cys Ala Ser Leu
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Lys Gly Pro Leu Gly Leu Lys Trp His Gln Met Val Asn Asn Ile Cys
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Gln Phe Ile Phe Trp Thr Gly Phe Ile Leu Met Leu Val Phe Tyr Val
195 200 205
Val Ile Ala Lys Lys Val Tyr Asp Ser Tyr Arg Lys Ser Lys Ser Lys
210 215 220
Asp Arg Lys Asn Asn Lys Lys Leu Glu Gly Lys Val Phe Val Val Val
225 230 235 240
Ala Val Phe Phe Val Cys Phe Ala Pro Phe His Phe Ala Arg Val Pro
245 250 255
Tyr Thr His Ser Gln Thr Asn Asn Lys Thr Asp Cys Arg Leu Gln Asn
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Leu Ser His Ser Tyr Cys Leu His Gln Asp Val Met Lys Leu Ala Cys
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Asp Asp Ile Arg Val Asn Val Val Tyr Gly Leu Ile Val Ile Ile Ser
195 200 205

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Ala Ile Gly Leu Asp Ser Leu Leu Ile Ser Phe Ser Tyr Leu Leu Ile
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225 230 235 240

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260 265 270

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Ile Tyr Ile Val Arg Thr Glu His Ser Leu His Glu Pro Met Tyr Ile
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Phe Leu Cys Met Leu Ser Gly Ile Asp Ile Leu Ile Ser Thr Ser Ser
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Met Pro Lys Met Leu Ala Ile Phe Trp Phe Asn Ser Thr Thr Ile Gln
85 90 95

Phe Asp Ala Cys Leu Leu Gln Met Phe Ala Ile His Ser Leu Ser Gly
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Met Glu Ser Thr Val Leu Leu Ala Met Ala Phe Asp Arg Tyr Val Ala
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Ile Cys His Pro Leu Arg His Ala Thr Val Leu Thr Leu Pro Arg Val
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Thr Lys Ile Gly Val Ala Ala Val Val Arg Gly Ala Ala Leu Met Ala
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Pro Leu Pro Val Phe Ile Lys Gln Leu Pro Phe Cys Arg Ser Asn Ile
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Leu Ser His Ser Tyr Cys Leu His Gln Asp Val Met Lys Leu Ala Cys
180 185 190

Asp Asp Ile Arg Val Asn Val Val Tyr Gly Leu Ile Val Ile Ile Ser
195 200 205

Ala Ile Gly Leu Asp Ser Leu Leu Ile Ser Phe Ser Tyr Leu Leu Ile
210 215 220

Leu Lys Thr Val Leu Gly Leu Thr Arg Glu Ala Gln Ala Lys Ala Phe
225 230 235 240

Gly Thr Cys Val Ser His Val Cys Ala Val Phe Ile Phe Tyr Val Pro
245 250 255

Phe Ile Gly Leu Ser Met Val His Arg Phe Ser Lys Arg Arg Asp Ser
260 265 270

Pro Leu Pro Val Ile Leu Ala Asn Ile Tyr Leu Leu Val Pro Pro Val
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Leu Thr Gln Ser Gln Asp Ile Gln Leu Leu Val Phe Val Leu Val Leu
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ata ttc tac ttc atc atc ctc cct gga aat ttt ctc att att ttc acc 144
Ile Phe Tyr Phe Ile Ile Leu Pro Gly Asn Phe Leu Ile Ile Phe Thr
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ata aag tca gac cct ggg ctc aca gcc ccc ctc tat ttc ttt ctg ggc 192
Ile Lys Ser Asp Pro Gly Leu Thr Ala Pro Leu Tyr Phe Phe Leu Gly
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atg Met 145	ttg Leu	gct Ala	ctg Leu	tgg Trp	ctt Leu 150	ggg Gly	ggt Gly	ttt Phe	gtc Val	cac His 155	tcc Ser	att Ile	atc Ile	cag Gln	gtg Val 160	480
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Ile Lys Ser Asp Pro Gly Leu Thr Ala Pro Leu Tyr Phe Phe Leu Gly
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Cys Ile Thr Gln Leu Phe Phe Leu His Phe Leu Gly Gly Gly Glu Gly
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145 150 155 160
Val Leu Ile Leu Arg Leu Pro Phe Cys Gly Pro Asn Gln Leu Asp Asn
165 170 175
Phe Phe Cys Asp Val Pro Gln Val Ile Lys Leu Ala Cys Thr Asp Thr
180 185 190
Phe Val Val Glu Leu Leu Met Val Phe Asn Ser Gly Leu Met Thr Leu
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Leu Cys Phe Leu Gly Leu Leu Ala Ser Tyr Ala Val Ile Leu Cys Arg
210 215 220
Ile Arg Gly Ser Ser Ser Glu Ala Lys Asn Lys Ala Met Ser Thr Cys
225 230 235 240
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245 250 255
Ile Tyr Thr Arg Pro Phe Arg Ala Phe Pro Ala Asp Lys Val Val Ser
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Ser Leu Gly Phe Ile Ile Gly Val Ser Val Val Gly Asn Leu Leu Ile
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Ser Ile Leu Leu Val Lys Asp Lys Thr Leu His Arg Ala Pro Tyr Tyr
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Gly Phe Met Leu Leu Leu Ala Leu Ile Leu Leu Ala Thr Gln Leu Val	
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Tyr Leu Lys Leu Ile Phe Phe Val His Asp Arg Arg Lys Met Lys Pro	
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Arg Ile Ser Arg Met Phe Tyr Ile Met Thr Phe Leu Phe Leu Thr Leu	
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Trp Gly Pro Tyr Leu Val Ala Cys Tyr Trp Arg Val Phe Ala Arg Gly	
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Pro Val Val Pro Gly Gly Phe Leu Thr Ala Ala Val Trp Met Ser Phe	
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Leu Arg Arg Cys Phe Ser Thr Thr Leu Leu Tyr Cys Arg Lys Ser Arg
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Val Ser Val Val Gly Asn Leu Leu Ile Ser Ile Leu Leu Val Lys Asp
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Lys Thr Leu His Arg Ala Pro Tyr Tyr Phe Leu Leu Asp Leu Cys Cys
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Ser Asp Ile Leu Arg Ser Ala Ile Cys Phe Pro Phe Val Phe Asn Ser
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Val Lys Asn Gly Ser Thr Trp Thr Tyr Gly Thr Leu Thr Cys Lys Val
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Ile Ala Phe Leu Gly Val Leu Ser Cys Phe His Thr Ala Phe Met Leu
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Phe Cys Ile Ser Val Thr Arg Tyr Leu Ala Ile Ala His His Arg Phe
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Tyr Thr Lys Arg Leu Thr Phe Trp Thr Cys Leu Ala Val Ile Cys Met
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Val Trp Thr Leu Ser Val Ala Met Ala Phe Pro Pro Val Leu Asp Val
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Gly Thr Tyr Ser Phe Ile Arg Glu Glu Asp Gln Cys Thr Phe Gln His
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Ser Leu Ala Ile Leu Ser Thr Glu Val Ser Val Leu Leu Leu Thr Phe	
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Thr Gly Phe Ile Val Ala Phe Ile Pro Leu Ser Asn Lys Glu Phe Phe	
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Gly Ile Asn Leu Ala Ala Phe Ile Ile Ile Val Phe Ser Tyr Gly Ser	
225 230 235 240	
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Val Ile Phe Ile Leu Pro Ile Asn Ser Ala Leu Asn Pro Ile Leu Tyr	
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Pro Glu Leu Met Lys Pro Asp Leu Phe Thr Tyr Pro Cys Glu Met Ser
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Val Phe Val Trp Val Val Ser Ala Val Thr Cys Phe Gly Asn Ile Phe
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Met Ser Ile Ile Ser Leu Cys Cys Ala Asp Cys Leu Met Gly Ile Tyr
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Leu Phe Val Ile Gly Gly Phe Asp Leu Lys Phe Arg Gly Glu Tyr Asn
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Lys His Ala Gln Leu Trp Met Glu Ser Thr His Cys Gln Leu Val Gly
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Ser Leu Ala Ile Leu Ser Thr Glu Val Ser Val Leu Leu Leu Thr Phe
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Leu Thr Leu Glu Lys Tyr Ile Cys Ile Val Tyr Pro Phe Arg Cys Val
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Arg Pro Gly Lys Cys Arg Thr Ile Thr Val Leu Ile Leu Ile Trp Ile
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Thr Gly Phe Ile Val Ala Phe Ile Pro Leu Ser Asn Lys Glu Phe Phe
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Asp Thr Glu Ser Ile Gly Ala Gln Ile Tyr Ser Val Ala Ile Phe Leu
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Gly Ile Asn Leu Ala Ala Phe Ile Ile Ile Val Phe Ser Tyr Gly Ser
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Asn Gln Val Lys Lys Glu Met Ile Leu Ala Lys Arg Phe Phe Phe Ile
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Val Phe Thr Asp Ala Leu Cys Trp Ile Pro Ile Phe Val Val Lys Phe
275 280 285

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Val Ile Phe Ile Leu Pro Ile Asn Ser Ala Leu Asn Pro Ile Leu Tyr
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325 330 335

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Pro Gly Lys Asn Thr Thr Leu His Asn Glu Phe Asp Thr Ile Val Leu
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gaa Glu	tct Ser	gca Ala	caa Gln	aaa Lys 290	atc Ile	cta Leu	tat Tyr	tac Tyr	tgc Cys 295	aaa Lys	gaa Glu	att Ile	aca Thr	ctt Leu 300	ttc Phe	915
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 His Arg Leu His Leu Val Val Tyr Ser Leu Val Leu Ala Ala Gly Leu
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Met Leu Ala Asn Ser Ser Ser Thr Asn Ser Ser Val Leu Pro Cys Pro
1 5 10 15

Asp Tyr Arg Pro Thr His Arg Leu His Leu Val Val Tyr Ser Leu Val
20 25 30

Leu Ala Ala Gly Leu Pro Leu Asn Ala Leu Ala Leu Trp Val Phe Leu
35 40 45

Arg Ala Leu Arg Val His Ser Val Val Ser Val Tyr Met Cys Asn Leu
50 55 60

Ala Ala Ser Asp Leu Leu Phe Thr Leu Ser Leu Pro Val Arg Leu Ser
65 70 75 80

Tyr	Tyr	Ala	Leu	His	His	Trp	Pro	Phe	Pro	Asp	Leu	Leu	Cys	Gln	Thr	
				85					90					95		
Thr	Gly	Ala	Ile	Phe	Gln	Met	Asn	Met	Tyr	Gly	Ser	Cys	Ile	Phe	Leu	
			100					105					110			
Met	Leu	Ile	Asn	Val	Asp	Arg	Tyr	Ala	Ala	Ile	Val	His	Pro	Leu	Arg	
		115					120					125				
Leu	Arg	His	Leu	Arg	Arg	Pro	Arg	Val	Ala	Arg	Leu	Leu	Cys	Leu	Gly	
	130					135					140					
Val	Trp	Ala	Leu	Ile	Leu	Val	Phe	Ala	Val	Pro	Ala	Ala	Arg	Val	His	
145					150					155					160	
Arg	Pro	Ser	Arg	Cys	Arg	Tyr	Arg	Asp	Leu	Glu	Val	Arg	Leu	Cys	Phe	
				165					170					175		
Glu	Ser	Phe	Ser	Asp	Glu	Leu	Trp	Lys	Gly	Arg	Leu	Leu	Pro	Leu	Val	
			180					185					190			
Leu	Leu	Ala	Glu	Ala	Leu	Gly	Phe	Leu	Leu	Pro	Leu	Ala	Ala	Val	Val	
		195					200					205				
Tyr	Ser	Ser	Gly	Arg	Val	Phe	Trp	Thr	Leu	Ala	Arg	Pro	Asp	Ala	Thr	
	210					215					220					
Gln	Ser	Gln	Arg	Arg	Arg	Lys	Thr	Val	Arg	Leu	Leu	Leu	Ala	Asn	Leu	
225					230					235					240	
Val	Ile	Phe	Leu	Leu	Cys	Phe	Val	Pro	Tyr	Asn	Ser	Thr	Leu	Ala	Val	
				245					250					255		
Tyr	Gly	Leu	Leu	Arg	Ser	Lys	Leu	Val	Ala	Ala	Ser	Val	Pro	Ala	Arg	
		260						265					270			
Asp	Arg	Val	Arg	Gly	Val	Leu	Met	Val	Met	Val	Leu	Leu	Ala	Gly	Ala	
		275					280					285				
Asn	Cys	Val	Leu	Asp	Pro	Leu	Val	Tyr	Tyr	Phe	Ser	Ala	Glu	Gly	Phe	
	290					295					300					
Arg	Asn	Thr	Leu	Arg	Gly	Leu	Gly	Thr	Pro	His	Arg	Ala	Arg	Thr	Ser	
305					310					315					320	
Ala	Thr	Asn	Gly	Thr	Arg	Ala	Ala	Leu	Ala	Gln	Ser	Glu	Arg	Ser	Ala	
				325					330					335		
Val	Thr	Thr	Asp	Ala	Thr	Arg	Pro	Asp	Ala	Ala	Ser	Gln	Gly	Leu	Leu	
			340					345					350			
Arg	Pro	Ser	Asp	Ser	His	Ser	Leu	Ser	Ser	Phe	Thr	Gln	Cys	Pro	Gln	
		355					360					365				
Asp	Ser	Ala	Leu													
			370													

<210> 21
 <211> 19
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<223> Description of Artificial Sequence: Primer LW1282

<400> 21
taatacctgc actgcccac

19

<210> 22
<211> 22
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer LW 1283

<400> 22
tctttccttc tcttctcact cc

22

<210> 23
<211> 32
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer LW 1373

<400> 23
gcataagctt atgctaacac tgaataaaaac ag

32

<210> 24
<211> 30
<212> DNA
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<220>
<223> Description of Artificial Sequence: Primer LW1374

<400> 24
gcattctcgag tcacatgctg taggatttgg

30

<210> 25
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<212> PRT
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<220>
<223> Description of Artificial Sequence: Peptide

<400> 25
Ala Pro Arg Thr Pro Gly Gly Arg Arg
1 5

<210> 26
<211> 32
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer LW1248

002227 01060300

<400> 26
gcatgaattc caatatactt ccccatacct ac 32

<210> 27
<211> 30
<212> DNA
<213> Artificial Sequence
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<223> Description of Artificial Sequence: Primer LW1249

<400> 27
gcatggatcc ggaaaagaag gagaagaaag 30

<210> 28
<211> 18
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer LW1278

<400> 28
accgctgcct ttttagtc 18

<210> 29
<211> 23
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer LW1279

<400> 29
ccttctttct gggtacataa gtc 23

<210> 30
<211> 30
<212> DNA
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<220>
<223> Description of Artificial Sequence: Primer LW1405

<400> 30
aagcataaca tggatgaaac aggaaatctg 30

<210> 31
<211> 29
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer LW1406

<400> 31
aagcataact atactttaca tatttcttc 29

<210> 32
<211> 22
<212> DNA
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<220>
<223> Description of Artificial Sequence: Primer LW1280

<400> 32
tctgcacaca gctcttccat gg 22

<210> 33
<211> 22
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer LW1281

<400> 33
tcccttggtcc agttggtga gg 22

<210> 34
<211> 30
<212> DNA
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<220>
<223> Description of Artificial Sequence: Primer LW1385

<400> 34
gcataagctt ccatggaact tcataacctg 30

<210> 35
<211> 30
<212> DNA
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<220>
<223> Description of Artificial Sequence: Primer LW1386

<400> 35
gcattctgag ttaccccccac agcgtgcag 30

<210> 36
<211> 29
<212> DNA
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<220>
<223> Description of Artificial Sequence: Primer LW1329

<400> 36
gcattctgag tcagcctaag gttatggtg 29

<210> 37
<211> 29
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<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Primer LW1377

<400> 37

gcataagctt atgaacacca cagtgatgc

29

<210> 38

<211> 41

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Primer LW1387

<400> 38

gagaaatatt tttctaaaaa aacctgtttt tgcaaaaacg g

41

<210> 39

<211> 41

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Primer LW1388

<400> 39

ccgtttttgc aaaaacaggt ttttttagaa aaatatttct c

41

<210> 40

<211> 30

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Primer LW1314

<400> 40

gcatgaattc ccacattcat catctacctc

30

<210> 41

<211> 29

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Primer LW1315

<400> 41

gcatggatcc gaagaccaa aagaccag

29

<210> 42

<211> 30

<212> DNA

<213> Artificial Sequence

<223> Description of Artificial Sequence: Primer LW1326

gcatgaattc atgatggtgg atccaatgg

30

<211> 27.

<212> DNA

<213> Artificial Sequence

<223> Description of Artificial Sequence: Primer LW1327

gcatctcgag cctagggctc tgaagcg

27

<211> 42

<212> DNA

<213> Artificial Sequence

<223>. Description of Artificial Sequence: Primer LW1415

ccatgtatat atttctttgc atgctttcag gcattgacat cc

42

<211> 42

<212> DNA

<213> Artificial Sequence

<223> Description of Artificial Sequence: Primer LW1416

ggatgtcaat gcctgaaagc atgcaaagaa atatatacat gg

42

<211> 30

<212> DNA

<213> Artificial Sequence

<223> Description of Artificial Sequence: Primer LW1308

gcatgaattc actcacttct catctccttc

30

<211> 30

<212> DNA

<213> Artificial Sequence

<223> Description of Artificial Sequence: Primer LW1309

<400> 47

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<220>
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27

<220>
<223> Description of Artificial Sequence: Primer LW1325.

35

<223> Description of Artificial Sequence: Primer LW1306

· 29

<220>
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28.

<223> Description of Artificial Sequence: Primer GV599

30

<210> 53
<211> 22
<212> DNA
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<220>
<223> Description of Artificial Sequence: Primer GV600

<400> 53
ctgaaacagc gcctcagctc cc

22

<210> 54
<211> 27
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer LW1482

<400> 54
agctatggcg aactatagcc atgcagc

27

<210> 55
<211> 27
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer LW148

<400> 55
agtcctcata taacacagta aggttcc

27

<210> 56
<211> 28
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer LW1310

<400> 56
gcatgaattc gcagaagaag gctattgg

28

<210> 57
<211> 29
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer LW1311

<400> 57
gcatggatcc gcagtaaaga agggttgtg

29

<210> 58
<211> 19

<212> DNA
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<220>
<223> Description of Artificial Sequence: Primer LW1442

<400> 58
gccattctgt ccacagaag

19

<210> 59
<211> 19
<212> DNA
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<220>
<223> Description of Artificial Sequence: Primer LW1443

<400> 59
tcagttgctg ttatggcac

19

<210> 60
<211> 24
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer LW1440

<400> 60
aagcggatgt ttagacctct tgtg
24

<210> 61
<211> 23
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer LW1441

<400> 61
aacagtcatt aataggaatt gag

23

<210> 62
<211> 32
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer LW1472

<400> 62
gcatgaattc tgccatgtca atcatttctc tc

32

<210> 63
<211> 31

$\langle 220 \rangle$

<400> 63

31

<210> 64

<211> 29

<212> DNA

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<400> 64

29

<210> 65

<211> 29

<212> DNA

<220>

<400> 65

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<210> 66

<211> 30

<212> DNA

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<211> 29

<212> DNA

<220>

<400> 67

29

<210> 68

<211> 43

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Primer 217A

<400> 68

taggtcggta gtcaggacac gggagaacag aactgttggg tga

43

<210> 69

<211> 52

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Primer 217B

<400> 69

gccccgtgtgg cggtttagat ccagaatgcc cattttctgt tccatctaac ca

52

<210> 70

<211> 20

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Primer LW1480

<400> 70

ggttctacct ggacttatgg

20

<210> 71

<211> 20

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Primer LW1481

<400> 71

taatgaatga gtaagtgcc

20

<210> 72

<211> 42

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Primer CON103a

<400> 72

tttattaata ttggaaggga caaactggag agcacagaac at

42

<210> 73

<211> 44

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Primer CON103b

<400> 73

44

<211> 18

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Primer 1332

tcctactgtc atgaaccc

18

<211> 18

<212> DNA

<213> Artificial Sequence

<22.0>

<223> Description of Artificial Sequence: Primer 1333

cagaagaagt tgtccagc

18

[illegible]